Zhou

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RAW SEQUENCE LISTING DATE: 05/24/2001
PATENT APPLICATION: US/09/601,561 TIME: 13:38:27

Input Set : A:\Nih332-1.app
Output Set: C:\CRF3\05242001\I601561.raw

ENTERED

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  3 <110> APPLICANT: Schneider, Thomas D.
  4       Hengen, Paul N.
  5       The Government of the United States of America
  6       as represented by The Secretary of the
  7       Department of Health and Human Services
  9 <120> TITLE OF INVENTION: Molecular Computing Elements: Gates and Flip-Flops
 11 <130> FILE REFERENCE: 015280-332100US
 13 <140> CURRENT APPLICATION NUMBER: US 09/601,561
C--> 14 <141> CURRENT FILING DATE: 2001-05-14
 16 <150> PRIOR APPLICATION NUMBER: US 60/075,468
 17 <151> PRIOR FILING DATE: 1998-02-20
 19 <150> PRIOR APPLICATION NUMBER: WO PCT/US99/03469
 20 <151> PRIOR FILING DATE: 1999-02-17
 22 <160> NUMBER OF SEQ ID NOS: 19
 24 <170> SOFTWARE: PatentIn Ver. 2.1
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 27 <211> LENGTH: 20
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 29 <213> ORGANISM: Artificial Sequence
 31 <220> FEATURE:
 32 <223> OTHER INFORMATION: Description of Artificial Sequence:consensus
 33       sequence of early model of Factor for Inversion
 34       Stimulation (Fis) binding site
 36 <400> SEQUENCE: 1
 37 ttgstcaaaa tttgascaaa                                              20
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 41 <211> LENGTH: 42
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 45 <220> FEATURE:
 46 <223> OTHER INFORMATION: Description of Artificial Sequence:paired Factor
 47       for Inversion Stimulation (Fis) binding sites with
 48       11 bp spacing; overlap 11
 50 <400> SEQUENCE: 2
 51 tattctttgc tcaaaatttg atcaaatttt gagcaaagaa ta                     42
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 56 <212> TYPE: DNA
 57 <213> ORGANISM: Artificial Sequence
 59 <220> FEATURE:
 60 <223> OTHER INFORMATION: Description of Artificial Sequence:paired Factor
 61       for Inversion Stimulation (Fis) binding sites with
 62       7 bp spacing; overlap 7
 64 <400> SEQUENCE: 3
 65 aggcttttgc tcaaagttta aactttgagc aaaagcct                          38
 68 <210> SEQ ID NO: 4
 69 <211> LENGTH: 15
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70 <212> TYPE: DNA
71 <213> ORGANISM: Artificial Sequence
73 <220> FEATURE:
74 <223> OTHER INFORMATION: Description of Artificial Sequence:sequence logo
75 for Factor for Inversion Stimulation (Fis) binding
76 site
78 <400> SEQUENCE: 4
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82 <210> SEQ ID NO: 5
83 <211> LENGTH: 58
84 <212> TYPE: DNA
85 <213> ORGANISM: Artificial Sequence
87 <220> FEATURE:
88 <223> OTHER INFORMATION: Description of Artificial Sequence:Factor for
89 Inversion Stimulation (Fis) binding sites
90 separated by 23 bp; separated 23
92 <400> SEQUENCE: 5
93 ggaattcttt gctcaaaatt tgatcaggat cctgatcaaa ttttgagcaa agaattcc 58
96 <210> SEQ ID NO: 6
97 <211> LENGTH: 21
98 <212> TYPE: DNA
99 <213> ORGANISM: Artificial Sequence
101 <220> FEATURE:
102 <223> OTHER INFORMATION: Description of Artificial Sequence:18.1 bit Fis
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110 <211> LENGTH: 21
111 <212> TYPE: DNA
112 <213> ORGANISM: Artificial Sequence
114 <220> FEATURE:
115 <223> OTHER INFORMATION: Description of Artificial Sequence:18.1 bit Fis
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118 <400> SEQUENCE: 7
119 tttgatcaaa ttttgagcaa a 21
122 <210> SEQ ID NO: 8
123 <211> LENGTH: 21
124 <212> TYPE: DNA
125 <213> ORGANISM: Artificial Sequence
127 <220> FEATURE:
128 <223> OTHER INFORMATION: Description of Artificial Sequence:12.7 bit Fis
129 site
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132 tttgctcaaa gtttaaactt t 21
135 <210> SEQ ID NO: 9
136 <211> LENGTH: 21
137 <212> TYPE: DNA
138 <213> ORGANISM: Artificial Sequence

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actcaaaaac tgaac 15

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Input Set : A:\Nih332-1.app
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212 <210> SEQ ID NO: 15
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217 <220> FEATURE:
218 <223> OTHER INFORMATION: Description of Artificial Sequence:synthesized
219 single very long nucleic acid with hairpin loop
220 DNA
222 <400> SEQUENCE: 15
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224 ccgatcgctc gaggaattcg aacaactgtt gttcagtttt tgagtggatc ccg 113
227 <210> SEQ ID NO: 16
228 <211> LENGTH: 21
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: Description of Artificial Sequence:8.4 bit Fis
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241 <211> LENGTH: 21
242 <212> TYPE: DNA
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246 <223> OTHER INFORMATION: Description of Artificial Sequence:10.0 bit Fis
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254 <211> LENGTH: 21
255 <212> TYPE: DNA
256 <213> ORGANISM: Artificial Sequence
258 <220> FEATURE:
259 <223> OTHER INFORMATION: Description of Artificial Sequence:10.0 bit Fis
260 site
262 <400> SEQUENCE: 18
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266 <210> SEQ ID NO: 19
267 <211> LENGTH: 21
268 <212> TYPE: DNA
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271 <220> FEATURE:
272 <223> OTHER INFORMATION: Description of Artificial Sequence:8.4 bit Fis
273 site
275 <400> SEQUENCE: 19
276 gttgttcagt ttttgagtgg a 21

VERIFICATION SUMMARY
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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date